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| FORM PTO-1449 U.S. DEPARTMENT OF COMMERCE PATENT AND TRADEMARK OFFICE | | | ATTY DOCKET NO. TSRI 923.1 | | SERIAL NO. 10/534,766 | |
| INFORMATION DISCLOSURE STATEMENT BY APPLICANT | | | APPLICANT Bracey, et al. | | | |
| | | | FILING DATE 11/14/2003 | | GROUP Unassigned | |
| U.S. PATENT DOCUMENTS | | | | | | |
| EXAM. INITIALS | DOCUMENT NUMBER | DATE | NAME | CLASS | SUB- CLASS | FILING DATE |
| | 5,221,410 | 6/22/93 | Kushner et al. | X | | |
| | 6,251,931 | 6/26/01 | Boger et al. | | | |
| | 6,271,015 | 8/7/01 | Gilula et al. | | | |
| | 6,462,054 | 10/8/02 | Boger et al. | | | |

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| FOREIGN PATENT DOCUMENTS | | | | | | |
| EXAM. INITIALS | DOCUMENT NUMBER | DATE | COUNTRY | CLASS | SUB- CLASS | TRANSLATION YES NO |
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| OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages) | | | |
| 1 | Bracey et al., "Structural adaptations in a membrane enzyme that terminates endocannabinoid signaling." <i>Science</i> , 298, pp. 1793-1796 (2002). | | |
| 2 | Boger et al., "Exceptionally potent inhibitors of fatty acid amide hydrolase: The enzyme responsible for degradation of endogenous oleamide and anandamide." <i>Proc. Natl. Acad. Sci. U.S.A.</i> , 97, pp. 5044-5049, (2000). | | |
| 3 | Collaborative Computational Project, Number 4, "The CCP4 Suite: Programs for Protein Crystallography", <i>Acta Cryst.</i> , D50, pp. 760-763 (1994). | | |
| 4 | Cravatt et al., "Molecular characterization of an enzyme that degrades neuromodulatory fatty-acid amides." <i>Nature</i> , 384, pp. 83-87, (1996). | | |
| 5 | Cravatt et al., "Fatty acid amide hydrolase: an emerging therapeutic target in the endocannabinoid system." <i>Current Opinion in Chemical Biology</i> , 7, pp. 469-475, (2003). | | |
| 6 | Egertová et al., "A new perspective on cannabinoid signalling: complementary localization of fatty acid amide hydrolase and the CB1 receptor in rat brain." <i>Proc. R. Soc. Lond. B. Biol. Sci.</i> , 265, pp. 2081-2085, (1998). | | |
| 7 | Garavito et al., "Strategies for crystallizing membrane proteins." <i>Journal of Bioenergetics and Biomembranes</i> , 28, pp. 13-27, (1996). | | |
| 8 | Giang et al., "Molecular characterization of human and mouse fatty acid amide hydrolases." <i>Proc. Natl. Acad. Sci. U.S.A.</i> , 94, pp. 2238-2242, (1997). | | |
| 9 | Kraulis, "MOLSCRIPT: a program to produce both detailed and schematic plots of protein structures." <i>J. Appl. Cryst.</i> , 24, pp. 946-950, (1991). | | |
| 10 | Labahn et al., "An Alternative Mechanism for Amidase Signaling Enzymes", <i>J. Mol. Bio.</i> , 322, pp. 1053-1064, (2002). | | |
| 11 | Makino et al., "Automated flexible ligand docking method and its applications for database search." <i>J. Comput. Chem.</i> , 18, pp. 1812-1825, (1997). | | |
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| 13 | Merritt et al., "Raster3D: Photorealistic Molecular Graphics", <i>Methods in Enzymology</i> , 277, pp. 505-524, (1997). | | |
| EXAMINER | /David J. Steadman/ | DATE CONSIDERED | 10/13/2008 |

EXAMINER: Initial if citation considered, whether or not citation is in conformance with MPEP 609; Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to Applicant.

ALL REFERENCES CONSIDERED EXCEPT WHERE LINED THROUGH. /DJS/

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| 14 | Nicholls et al., "Protein Folding and Association: Insights From Interfacial and Thermodynamic Properties of Hydrocarbons", <i>Proteins Struct. Funct. Genet.</i> , 11, pp. 281-296, (1991). |
| 15 | Patricelli et al., "Comparative Characterization of a Wild Type and Transmembrane Domain-Deleted Fatty Acid Amide Hydrolase: Identification of the Transmembrane Domain as a Site for Oligomerization", <i>Biochemistry</i> , 37, pp. 15177-15187, (1998). |
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| 18 | Patricelli et al., "Characterization and Manipulation of the Acyl Chain Selectivity of Fatty Acid Amide Hydrolase", <i>Biochemistry</i> , 40, pp. 6107-6115, (2001). |
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| 20 | Rice et al., "Single-wavelength anomalous diffraction phasing revisited." <i>Biological Crystallography</i> , D56, pp. 1413-1420, (2000). |
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| 22 | Shin et al., "Structure of malonamidase E2 reveals a novel Ser-cisSer-Lys catalytic triad in a new serine hydrolase fold that is prevalent in nature." <i>Embo J.</i> , 21, pp. 2509-2516, (2002). |
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| 25 | Wendt et al., "Structure and Function of a Squalene Cyclase", <i>Science</i> , 277, pp. 1811-1815, (1997). |
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